

STEATOHEPATITIS

Rodent nutritional model of non-alcoholic steatohepatitis: Species, strain and sex difference studies

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Abstract

Background and Aim: The methionine choline-deficient (MCD) diet leads to steatohepatitis in rodents. The aim of the present study was to investigate species, strain and sex differences in this nutritional model of non-alcoholic steatohepatitis (NASH).

Methods: Male and female Wistar, Long-Evans and Sprague-Dawley rats, and C57/BL6 mice ($n = 6$ per group) were fed a MCD diet for 4 weeks. Control groups received an identical diet supplemented with choline bitartrate (0.2% w/w) and methionine (0.3% w/w). Liver pathology (steatosis and inflammation) and ultrastructure, liver lipid profile (total lipids, triglycerides, lipid peroxidation products), liver: body mass ratios and serum alanine aminotransferase (ALT) levels were compared between these groups.

Results: The MCD diet-fed male rats developed greater steatosis ($P < 0.001$), had higher liver lipid content ($P < 0.05$) and had higher serum ALT levels ($P < 0.005$) than did female rats. Wistar rats (both sexes) had higher liver lipid levels ($P < 0.05$), serum ALT levels ($P < 0.05$), and liver mass: body mass ratios ($P < 0.025$) than did Long-Evans and Sprague-Dawley rats. In female groups, Wistar rats showed greater fatty change than did the other two strains ($P < 0.05$). All rats fed the MCD diet developed hepatic steatosis, but necrosis and inflammation were minor features and fibrosis was absent. Compared with Wistar rats, male C57/BL6 mice showed a marked increase in inflammatory foci ($P < 0.001$), and products of lipid peroxidation (free thiobarbituric acid reactive substances) ($P < 0.005$), and mitochondrial injury, while showing less steatosis ($P < 0.005$), lower hepatic triglyceride levels, ($P < 0.005$) and lower early lipid peroxidation products (conjugated dienes and lipid hydroperoxides; $P < 0.005$ and $P < 0.01$, respectively).

Conclusions: The Wistar strain and the male sex are associated with the greatest degree of steatosis in rats subjected to the MCD diet. Of the groups studied, male C57/BL6 mice develop the most inflammation and necrosis, lipid peroxidation, and ultrastructural injury, and best approximate the histological features of NASH.

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Key words: choline, lipid, lipid peroxidation, methionine, mice, mitochondria, non-alcoholic steatohepatitis, rats, sex, strain.

INTRODUCTION

Non-alcoholic fatty liver disease is one of the common forms of liver disease seen in outpatient practice.^{1,2} It is associated with a wide spectrum of morphological manifestations, including fatty change (steatosis), inflammation, necrosis, and occasionally cirrhosis. Overlap

between these patterns of injury often exists, and steatosis is frequently associated with hepatocyte injury and inflammation; a condition known as non-alcoholic steatohepatitis (NASH).^{3,4}

Until recently, NASH was regarded as a mild disease that rarely progresses to cirrhosis.^{3,5,6} However, it is now recognized that a significant proportion of

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patients with NASH do develop progressive fibrosis leading to cirrhosis, and occasionally, hepatocellular carcinoma.^{1,4} Non-alcoholic steatohepatitis is seen in association with obesity, type 2 diabetes mellitus, hyperlipidemia, rapid weight loss and jejunoileal bypass surgery for obesity;¹⁰ a similar pattern of liver injury can occur following exposure to certain drugs and industrial toxins.¹¹

Animal models have greatly contributed to the understanding of NASH. While several models of steatosis exist,¹² fewer models of steatohepatitis are available. One such model is the high-fat methionine choline-deficient diet (MCD),^{13,14} which results in hepatic steatosis, necroinflammation, and, in the longer term, fibrosis.¹⁴ Although initially established as a model of fatty liver transplantation,¹⁵ its application to the study of NASH pathogenesis has provided important new insights.¹⁶⁻²⁰ Steatohepatitis in this model is associated with upregulated hepatocyte microsomal cytochrome P450 2E1 (CYP2E1) activity and increased hepatic lipid peroxidation.^{14,15} CYP4A proteins provide an alternative source of microsomal pro-oxidant activity in CYP2E1 nullizygous mice fed the MCD diet.¹⁸ The importance of these findings is highlighted by the subsequent demonstration of enhanced CYP2E1 expression,²¹ as well as by by-products of oxidative stress²² in liver biopsies from patients with NASH. Inflammation and fibrosis in the MCD model is leptin-dependent,¹⁶ suggesting that leptin may play a regulatory role in hepatic fibrogenesis in NASH. Recent studies suggest that peroxisome proliferation activator receptor (PPAR)- α -regulated pathways may protect against the development of steatohepatitis, because PPAR- α agonists can prevent steatosis and inflammation in animals fed a high-fat MCD diet.¹⁷

To date there have been no systematic studies that have investigated the influence of species, strain and sex differences on the degree of hepatic steatosis, inflammation and fibrosis in the MCD nutritional model of NASH. Previous studies have used male Wistar,^{14,18} ACT,¹³ and Sprague-Dawley rats,¹⁸ male C57/BL6 mice,^{15,17,20} and genetically obese male (ob/ob) mice.¹⁶ In humans, genetic factors²³ and sex differences²⁴⁻²⁷ are known to influence individual susceptibility to NASH and the female sex is an independent predictor of fibrotic severity in NASH.²⁸ The present study thus sought to further characterize this model by testing the hypothesis that species, strain and sex differences influence liver injury in the MCD nutritional model of NASH.

METHODS

The present study was approved by the Animal Research Review Committee of the Faculty of Medicine, University of Cape Town, South Africa.

Materials

The pelleted MCD diet and the MCD diet supplemented with choline bitartrate (2 g/kg) and DL-

methionine (3 g/kg) were custom-made by ICN Biomedicals (Irvine, CA, USA). Chloroform, methanol and glutaraldehyde were obtained from Merck (Darmstadt, Germany). Osmium tetroxide was from Next Chimica (Centurion, South Africa). Spurr's resin was obtained from Agar Scientific limited (Stanstead, UK). All other materials were obtained from Sigma (St Louis, MO, USA).

Animals and diet

Wistar rats were obtained from the Medical Research Council Animal Unit (Cape Town, South Africa), Long-Evans rats and C57/BL6 mice from the University of Cape Town Animal Unit, and Sprague-Dawley rats from the University of Witwatersrand Animal Unit (Johannesburg, South Africa). All animals were 12 weeks of age at the commencement of the study. Animals were allowed to acclimatize to their new conditions for 1 week prior to the commencement of the study. The eight study groups, comprising male and female rats of the three different rat strains and C57/BL6 mice ($n = 6$ per group) were fed the MCD diet for 4 weeks, while control groups ($n = 6$ per group) received an identical diet to which choline bitartrate (2 g/kg) and DL-methionine (3 g/kg) was added. An additional control group was included that received standard rodent food. Rats were given free access to food and water and were weighed at weekly intervals for the duration of the study. The duration of the study and the composition of the diets were based on those used in previous studies.¹⁴⁻²⁰

Tissue preparation and analysis

At the end of the study, animals were either anesthetized, killed, and their livers were removed and weighed and sampled for histological assessment. Additional samples were snap frozen in liquid nitrogen for later analysis of liver lipid levels and lipid peroxidation. Samples for electron microscopy were fixed in 2.5% glutaraldehyde in phosphate buffered saline (pH = 7.4).

Biochemical analyses

Scrum was assayed for alanine aminotransferase (ALT) activity spectrophotometrically as described by Harder and Rej.²⁹ Frozen liver samples were thawed on ice, homogenized and sonicated with chilled SEAP buffer (0.15 mol/L NaCl, 0.001 mol/L EDTA, 0.01 g% NaN₃, 0.002 mol/L NaH₂PO₄, pH = 7.4), and brought to a final volume of 2 mL, of which 800 μ L was used for lipid extraction (triglycerides, conjugated dienes and lipid hydroperoxides) and the remainder for other assays. Lipid extraction was performed according to Folch,³⁰ as modified by Bligh and Dyer,³⁰ and extracts were dried under nitrogen jets. Triglycerides were measured colorimetrically in lipid extracts using a commercial kit (Roche, Man-

nheim, Germany) as previously reported.³¹ Early, intermediate and late products of lipid peroxidation (conjugated dienes, lipid hydroperoxides and thiobarbituric acid reactive substances [TBARS], respectively) were measured by spectrophotometric assays. Conjugated dienes and lipid hydroperoxides were measured in liver lipid extracts as described by Vasankari *et al.*³² and Ziaang *et al.*³³, respectively. Free and total TBARS in liver homogenates were measured by the method described by Asakawa *et al.*³⁴ Assays were performed both in the presence and absence of the antioxidant butylated hydroxytoluene (BHT); BHT limits the generation of TBARS during the assay, ensuring that only the products present in the liver at the time of death are measured (free TBARS).³⁴ Protein concentrations in liver homogenates were measured using the Markwell modification of the Lowry assay.³⁵ Hepatic lipid indices were expressed per milligram of protein.

Histopathology

Sections of formalin-fixed, paraffin-embedded samples were stained with H&E and analyzed using light microscopy. Selected samples were stained with Sirius red. Fatty change was graded according to the percentage of hepatocytes containing macrovesicular fat (grade 1: 0–25%; grade 2: 26–50%; grade 3: 51–75%; grade 4: 76–100%). Necroinflammation was quantified histologically by counting inflammatory foci, arbitrarily defined as groups of five or more leukocytes in 20 consecutive high-power fields (×40 objective). Assessment of liver injury was performed using coded slides to avoid observer bias.

Electron microscopy

For ultrastructural analysis, 1 mm³ samples were fixed in glutaraldehyde (2.5%), postfixed in osmium tetroxide

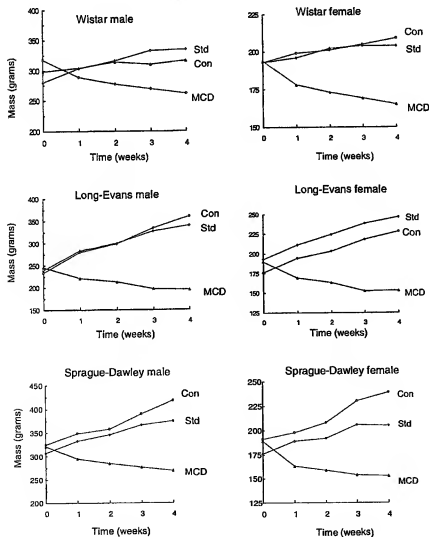


Figure 1 Body masses of rats fed methionine choline-deficient (MCD) diet, control (MCD supplemented with choline [2 g/kg] and methionine [3 g/kg]), and standard diets. Data are expressed as the mean masses of the six animals in each group.

ide (2%) and embedded in Spurr's resin. Sections were stained with uranyl acetate (2%, w/v) and lead citrate prior to analysis with electron microscopy (Zeiss EM 109; Oberkochen, Germany). For quantification of mitochondrial size, five random fields (magnification $\times 20\,000$) from five random cells in each section were photographed by an electron microscopist blinded to treatment groups; each photograph contained 5–10 mitochondria in a particular cell. Morphometric measurement of the mitochondrial area was performed on the five largest mitochondria per photograph using the JLGEnias image analysis software (Joyce-Loebl, Gateshead, UK).

Statistical analysis

Data was subjected to statistical analysis using the Kruskal-Wallis test and the Student's *t*-test where applicable.

RESULTS

Rat study: Strain and sex differences

Animal growth

In all groups, animals fed the MCD diet lost weight compared with those fed control or standard rodent diets; the latter showed a steady increase in weight (Fig. 1). Apart from weight loss, the general condition of the animals remained satisfactory.

Liver mass and morphology

The livers of male and female Wistar rats fed the MCD diet increased in mass relative to controls, while those from male and female Long-Evans and Sprague-Dawley rats decreased in mass (Table 1). To control for the loss of body mass induced by the MCD diet, liver mass was expressed as a percentage of body mass (relative liver mass). Male and female Wistar rats fed the MCD diet showed marked increases in relative liver mass while minimal changes were seen in MCD-fed Long-Evans and Sprague-Dawley rats ($P < 0.0025$). Macroscopically, livers from Wistar rats fed the MCD diet appeared pale and fatty compared with controls; these changes were less noticeable in MCD-diet-fed Long-Evans and Sprague-Dawley rats (Fig. 2).

Serum alanine aminotransferase levels

Serum ALT levels were significantly higher in all groups fed the MCD diet compared with controls ($P < 0.05$), except in the case of female Sprague-Dawley rats (Fig. 3). Wistar rats fed the MCD diet (male and female) had higher ALT levels than did their counterparts from other strains ($P < 0.01$ and $P < 0.05$, respectively). As a group, MCD-diet-fed male rats had higher ALT levels than their female counterparts ($P < 0.02$).

With respect to liver weights, liver histology and serum transaminase levels, there were no differences

Table 1 Liver mass (absolute and relative) and changes in relative liver mass resulting from the methionine choline-deficient (MCD) diet (relative liver mass [MCD]/relative liver mass [control] $\times 100$) in the six study groups

	MCD						Control					
	Wistar		Long-Evans		Sprague-Dawley		Wistar		Long-Evans		Sprague-Dawley	
	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female	Male	Female
Liver mass (grams)	12.50 \pm 3.16	8.63 \pm 0.62	7.88 \pm 1.64	4.95 \pm 1.28	10.08 \pm 1.10	4.87 \pm 0.42	10.40 \pm 2.51	6.42 \pm 0.55	13.00 \pm 1.64	7.40 \pm 0.77	14.10 \pm 1.79	7.70 \pm 0.65
Relative liver mass (liver mass/body mass) $\times 100$	4.73 \pm 0.97	5.23 \pm 0.28	4.09 \pm 0.75	3.20 \pm 0.68	3.72 \pm 0.26	3.20 \pm 0.30	3.27 \pm 0.21	3.05 \pm 0.22	3.60 \pm 0.42	3.30 \pm 0.55	3.30 \pm 0.45	3.22 \pm 0.22
Percentage change in relative liver mass	45	72	12	0	12	0	NA	NA	NA	NA	NA	NA

Relative liver mass = liver mass as a percentage of body mass. Data are expressed as mean \pm SD for each group. NA, not applicable.

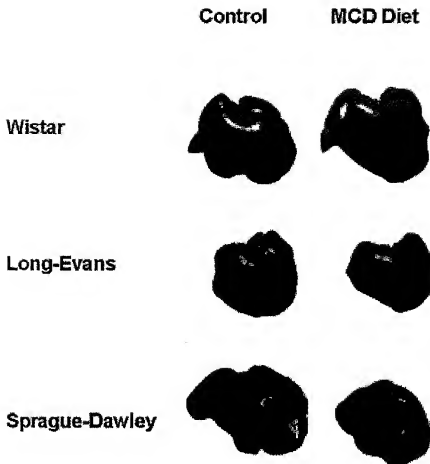


Figure 2 Macroscopic appearance of livers from male Wistar, Long-Evans and Sprague-Dawley rats fed control or methionine choline-deficient (MCD) diets. Female counterparts showed similar macroscopic features.

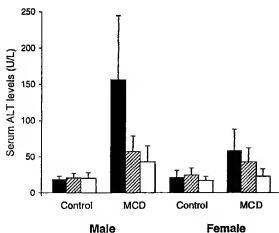


Figure 3 Serum alanine aminotransferase (ALT) levels in male and female (■) Wistar, (▨) Long-Evans and (□) Sprague-Dawley rats fed methionine choline-deficient (MCD) or control diets for 4 weeks. Data are expressed as mean \pm SD for each group.

between animals fed the control diet (MCD diet plus choline bitartrate [2 g/kg] and DL-methionine [3 g/kg]) and those receiving standard rodent diet (data not shown). Therefore, only the data for the group receiving the control diet is presented.

Liver lipid content

The liver lipid content (expressed as mg lipid/100 mg liver weight) was higher in male and female Wistar rats fed the MCD diet than in their counterparts in the other two strains ($P < 0.05$ and $P < 0.02$, respectively) (Fig. 4). As a group, male rats had higher liver lipid levels than female rats ($P < 0.002$).

Liver pathology

Livers from animals fed the MCD diet showed macrovesicular steatosis that varied in quantity and distribution in the various groups (Fig. 5). In the least affected groups (female Long-Evans and Sprague-Dawley rats) steatosis was confined to zone 3, while in the most severely affected animals (male Wistar rats)

close to 100% of hepatocytes showed macrovesicular steatosis. When steatosis was graded according to the percentage of hepatocytes containing macrovesicular fat, Wistar rats had higher scores than did the other

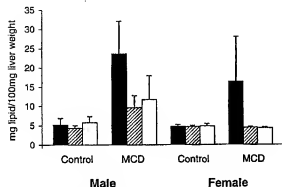


Figure 4 Total liver lipid content (mg/100 mg liver) in male and female (■) Wistar, (▨) Long-Evans and (□) Sprague-Dawley rats fed methionine choline-deficient (MCD) or control diets for 4 weeks. Data are expressed as mean \pm SD for each group.

strains, although this was only statistically significant in female rats ($P < 0.05$) (Fig. 6). As a group, male rats showed a greater degree of steatosis than did female rats ($P < 0.0002$). In all rats fed the MCD diet, inflammation and necrosis were minor features, while fibrosis was absent.

Mouse study: Comparison of C57/BL6 mice and Wistar rats

Male and female C57/BL6 mice fed the MCD diet were analyzed with respect to serum ALT levels, liver histology (steatosis and necroinflammation), ultrastructure and lipid biochemistry and compared with Wistar rats, the most susceptible rat strain.

Male C57/BL6 mice showed a marked increase in necroinflammatory foci compared with female C57/BL6 mice ($P < 0.02$) and Wistar rats of either sex ($P < 0.001$) (Table 2, Fig. 7a,b). These necroinflammatory foci were composed predominantly of lymphocytes with occasional neutrophils. Hepatocyte ballooning and Mallory bodies were not present. In some mice, focal pericellular fibrosis was noted (Fig. 7c). Both male and female mice showed less steatosis than did the Wistar

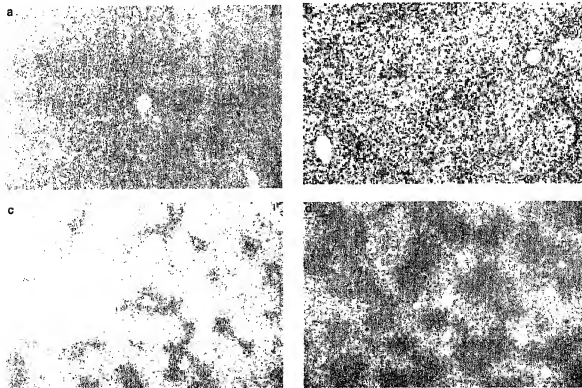


Figure 5 HE-stained sections from rats fed the methionine choline-deficient (MCD) or control diets for 4 weeks, illustrating differences in fatty change. (a) Control rat (magnification $\times 40$). The histological appearance is normal and representative of controls from all groups. (b) A male Wistar rat fed the MCD diet. Fatty change involves the entire liver lobule. (c) Female Wistar rat fed the MCD diet. Fatty change involves zones 3 and 2 with periportal sparing. (d) Female Long-Evans rat fed the MCD diet. Fatty change is largely confined to zone 3.

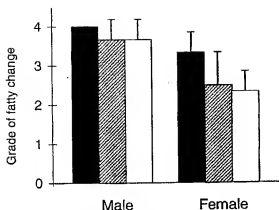


Figure 6 Average histological grade of fatty change in male and female (■) Wistar, (▨) Long-Evans and (□) Sprague-Dawley rats fed the methionine choline-deficient diet for 4 weeks. Fatty change was graded histologically according to the percentage of hepatocytes containing macrovesicular fat (grade 1: 0–25%; grade 2: 26–50%; grade 3: 51–75%; grade 4, 76–100%). Data are expressed as mean \pm SD for each group.

rats ($P < 0.005$). Serum ALT levels were elevated in male and female MCD-fed mice compared with controls ($P < 0.005$); however, the differences between male and female rodents and between C57/BL6 mice and Wistar rats were not statistically significant (Table 2).

Male C57/BL6 mice and Wistar rats were further studied with respect to hepatic ultrastructural morphology and lipid peroxidation; female rodents were not included in these analyses because of the minor degree of liver injury seen in these animals.

Ultrastructural morphology

Male C57/BL6 mice and Wistar rats fed the MCD diet showed evidence of ultrastructural injury that was not observed in controls; this was more pronounced in C57/BL6 mice. Such changes included profound mitochondrial swelling, proliferation of the smooth endoplasmic reticulum, loss of the normal relationship between the rough endoplasmic reticulum and mitochondria, and blebbing of the nuclear membrane (Fig. 7d and 8). In MCD-diet-fed mice, most cells contained a number of enlarged mitochondria, some of which were 3–4-fold the normal size; no such mitochondria were noted in control animals. Morphometric analysis of confirmed differences in mitochondrial size in MCD-diet-fed and control mice (average mitochondrial surface area: control $0.38 \pm 0.12 \mu\text{m}^2$ vs MCD $0.71 \pm 0.16 \mu\text{m}^2$; $P < 0.03$). Paracrystalline inclusions, reported in the mitochondria of humans²⁹ and animals³⁰ with NASH, were not seen in the present study.

Male C57/BL6 mice and Wistar rats were also compared with respect to hepatic lipid biochemistry. Triglyceride levels were far higher in MCD-diet-fed Wistar rats compared with their murine counterparts ($P < 0.005$) (Table 3). The MCD-diet-fed rodents

Table 2 Average histological grades of fatty change, inflammatory activity, and serum alanine aminotransferase (ALT) levels in male and female Wistar and C57/BL6 mice fed methionine choline-deficient (MCD) and control diet for 4 weeks

	MCD				Control			
	Wistar		C57/BL6		Wistar		C57/BL6	
	Male	Female	Male	Female	Male	Female	Male	Female
Inflammatory foci per 20 high-power field	5.00 \pm 3.29	6.50 \pm 5.13	41.67 \pm 25.38	6.67 \pm 0.82	0.67 \pm 0.82	0.17 \pm 0.41	2.75 \pm 1.26	2.33 \pm 1.97
Fatty change (average histological grade)	4.00 \pm 0.00	3.3 \pm 0.52	2.67 \pm 0.52	2.33 \pm 0.82	NA	NA	NA	NA
Serum ALT levels (IU/L)	136.67 \pm 29.62	58.67 \pm 29.62	99.83 \pm 36.76	70.67 \pm 16.17	18.83 \pm 4.62	21.50 \pm 9.71	18.67 \pm 6.77	21.33 \pm 12.23

^aInflammatory focus defined arbitrarily as a collection of ≥ 5 leukocytes. Data are expressed as mean \pm SD for each group. NA, not applicable.

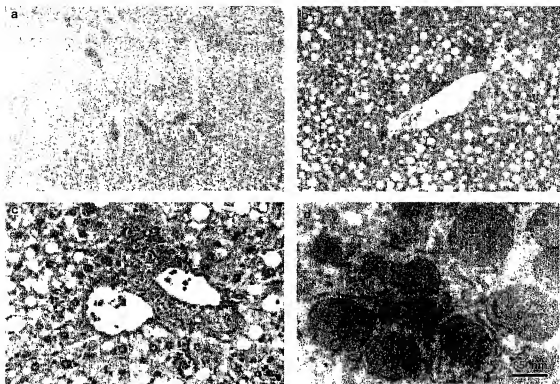


Figure 7 (a,b) HE-stained sections from the liver of a male C57/BL6 mouse fed the methionine choline-deficient (MCD) diet for 4 weeks ($\times 40$ and $\times 200$ magnification, respectively) illustrating mild steatosis but extensive necroinflammation. (c) Bile Sirius red stained sections illustrating pericellular fibrosis in an MCD-diet-fed male C57/BL6 mouse. Collagen is stained red. (d) Representative transmission electron micrograph illustrating mitochondrial enlargement in a male C57/BL6 mouse fed the MCD diet (compare with control mitochondria, Fig. 8a).

showed elevated levels of early, intermediate and late products of lipid peroxidation compared with their respective controls (Table 3). The MCD-diet-fed rats had higher levels of conjugated dienes and lipid hydroperoxides (early and intermediate products of lipid peroxidation) than did mice ($P < 0.005$ and $P < 0.01$, respectively); in contrast, mice showed far greater levels of free TBARS (end-products of lipid peroxidation) than did rats ($P < 0.005$). Total TBARS, which reflect merely the total available peroxidizable lipid, were more elevated in rats, in keeping with elevated triglyceride levels in these animals.

DISCUSSION

The present study has identified species, strain and sex differences in the MCD rodent nutritional model of NASH. Among the rat strains studied, male Wistars developed the greatest degree of steatosis when fed the MCD diet; in all rats inflammation and necrosis were minor features and fibrosis was absent. In contrast, male C57/BL6 mice fed the MCD diet showed florid hepatic necroinflammation with a lesser degree of steatosis. Consistent with previous

studies, all animals fed the MCD diet lost weight, while controls showed a steady increase in body weight.^{13,14}

The increased susceptibility of male rodents to liver injury in this nutritional model of NASH differs from human NASH, in which women show both a higher prevalence of NASH and increased fibrotic activity.³²⁻³⁷ Biochemical mechanisms unique to the MCD model may be responsible for the sex differences observed in the present study. The proposed biochemical basis for steatosis in this model is a block in phosphatidylcholine (PC) synthesis resulting from a dietary deficiency in methionine and choline.³⁷ In animals and in humans, PC synthesis may occur via two pathways: one that involves the direct incorporation of preformed choline into phosphatidyl compounds, and the other that is dependent upon the stepwise methylation of phosphatidylethanolamine by *S*-adenosyl methionine.³⁸⁻⁴⁰ Both pathways of PC synthesis are thus blocked by a dietary deficiency in choline and methionine. Because PC is an essential component of very-low-density lipoprotein, which is responsible for triglyceride clearance from hepatocytes, PC deficiency leads to the accumulation of triglycerides within hepatocytes, and thus steatosis.³⁷ Several studies have demonstrated sex differences in the

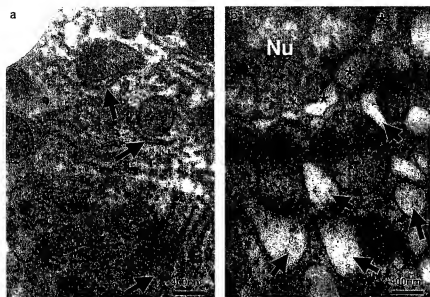


Figure 8 (a) Transmission electron micrograph of a representative hepatocyte from a control animal showing normal-sized mitochondria surrounded by irregularly disposed cysternae of rough endoplasmic reticulum (arrows). Smooth endoplasmic reticulum is not prominent. (b) Transmission electron micrograph of a representative hepatocyte from a mouse fed the MCD diet showing proliferation of smooth endoplasmic reticulum (arrows) and loss of the normal relationship between rough endoplasmic reticulum and mitochondria. Blebbing of the nuclear membrane (*) was frequently observed.

Table 3 Hepatic triglycerides and lipid peroxidation products including thiobarbituric acid reactive substances (TBARS, total and free), lipid hydroperoxides and conjugated dienes in male Wistar rats and C57/BL6 mice fed methionine choline-deficient (MCD) or control diets for 4 weeks. Data are expressed as mean \pm SD for each group

	Wistar rats		C57/BL6 mice	
	Control	MCD	Control	MCD
Triglycerides (μ g/mg protein)	92.7 \pm 13.4	558.0 \pm 140.0	71.6 \pm 18.0	172.0 \pm 12.9
TBARS (free) (nmol/mg protein)	0.44 \pm 0.12	2.20 \pm 1.30	1.30 \pm 2.60	11.20 \pm 2.70
TBARS (total) (nmol/mg protein)	4.5 \pm 0.7	307.1 \pm 29.1	5.2 \pm 1.7	81.9 \pm 25.1
Lipid hydroperoxides (nmol/mg protein)	0.53 \pm 0.16	8.70 \pm 0.70	1.20 \pm 1.20	6.60 \pm 1.30
Conjugated dienes (μ mol/mg protein)	7.11 \pm 0.27	27.60 \pm 5.50	7.70 \pm 3.00	12.12 \pm 3.60

pathways of PC synthesis between animals and humans, with methionine-dependent pathways apparently favored by estrogen.³⁹⁻⁴¹ Female rats fed a choline- (but not methionine-) deficient diet have been shown to develop far less hepatic steatosis than male rats.⁴⁰ This difference has been attributed to the greater propensity of females to use the methionine-dependent pathway of PC synthesis.⁴⁰ In the MCD model, however, where methionine deficiency is combined with choline deficiency, female rats would be expected to lose this apparent 'advantage', as they can no longer exploit the methionine-dependent pathway of PC synthesis. This was not observed in the present study where female rodents consistently developed less steatosis than male rodents. One possible explanation may be the ability of methionine to undergo regeneration following methylation reactions during PC synthesis. In such reactions, only the methyl group in S-adenosyl-methionine is consumed; the remainder of the molecule is intact as S-adenosylhomocysteine, which can be hydrolyzed to release free homocysteine. Subsequent methylation of homocysteine results in the regeneration of methionine,

with methyl groups being donated by cytosolic 5,10-methylene- H_2 -folate.³⁹ Thus, methionine-dependent pathways of PC synthesis may continue to operate, albeit at a lower level, in the presence of a dietary deficiency of methionine, allowing female rodents to retain their relative advantage over male rodents. Sex differences in metabolic pathways are well recognized in other rodent models of hepatic injury,⁴² and in the hepatic metabolism of drugs in animals and humans.^{43,44}

Marked strain-related differences in steatosis were also identified in rats fed the MCD, with Wistar rats showing the greatest propensity to develop steatosis. The mechanisms underlying these differences are uncertain. It is possible that subtle variations in PC synthesis, as described above, may exist, or that genetic differences at the level of hepatic free fatty acid (FFA) uptake, endogenous FFA synthesis, or β -oxidation of FFA may play a role. Strain-related differences in metabolic pathways and related liver injury have been demonstrated in other animal models of hepatotoxicity, such as the hypoxic halothane model.⁴⁵ Recent studies of ani-

mals with inherited mutations involving genes regulating hepatic carbohydrate and lipid metabolism, and in genetically manipulated mice, suggest that genetic factors may play a role in the pathogenesis of NASH.¹² While the influence of genetic factors in human NASH remains unclear, familial clustering of NASH in kindreds has recently been reported.³

The rodent group most profoundly affected by the MCD diet was the male C57/BL6 mice. In contrast to other groups, these animals developed florid necroinflammation and showed some evidence of early perivascular fibrosis. Interestingly, C57/BL6 mice developed less steatosis than did rats, suggesting that the increased necroinflammation is not a function of steatosis, but more likely of downstream events such as lipid peroxidation. Lipid peroxidation can injure hepatocytes both directly and indirectly through the pro-inflammatory and pro-fibrogenic properties of its end-products, malondialdehyde and 4-hydroxynonenal.¹⁴ Indeed, the present study found end-products of lipid peroxidation (free TBARS) to be markedly elevated in male C57/BL6 mice compared with male Wistar rats. Interestingly, early and intermediate products of lipid peroxidation (conjugated dienes and lipid peroxides, respectively) were higher in rats than in mice, and this may reflect the increased availability of lipid substrate in these animals and/or less progression to end-products. Mice, in contrast, have less steatosis and lower levels of early lipid peroxide products, but a greater propensity to take the lipid peroxidation process to completion (as evidenced by the higher free TBARS levels). The total TBARS merely reflect the total lipid available for lipid peroxidation *in vitro* and, in the present study, appear to correlate with the degree of steatosis and hepatic triglyceride levels.

Male C57/BL6 mice showed ultrastructural evidence of hepatocyte injury, including profound mitochondrial enlargement, blebbing of nuclear membranes, and dissociation of rough endoplasmic reticulum from mitochondria. The proliferation of smooth endoplasmic reticulum seen in these animals is consistent with the upregulation of CYP2E1 reported in this model.^{14,15} Although ultrastructural changes were also present in MCD-fed Wistar rats, these were less pronounced than in mice. Para-crystalline intramitochondrial inclusions that have been reported in human patients with NASH¹⁶ were not observed in this study; such inclusions have, however, been observed in animals fed the MCD diet for longer periods.²⁰ Mitochondrial injury in animals fed the MCD diet may be a result of oxidative stress and consequent lipid peroxidation of mitochondrial membranes. Mitochondrial injury activates hepatocyte apoptotic mechanisms and, when more severe, results in ATP depletion and hepatocyte necrosis,⁴⁶ a prominent feature in male C57/BL6 mice. The increased lipid peroxidation (evidenced by increased free TBARS) in the C57/BL6 mice compared with Wistar rats may, in part, explain differences in necroinflammation in these animals.

In conclusion, the present study has demonstrated profound species, strain and sex differences in the MCD nutritional model of NASH. Of the groups stud-

ied, male C57/BL6 mice develop the histological features that most closely resemble those seen in human NASH.

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Steatohepatitis: A Tale of Two "Hits"?

See article on page 764.

It has long been recognized that hepatic steatosis (fatty liver) occurs frequently in heavy alcohol drinkers and obese individuals.^{1,2} It may also follow the ingestion of a wide variety of therapeutic drugs.³ Steatosis of any etiology can be associated with the development of necroinflammation and fibrosis, so-called steatohepatitis, and even cirrhosis.⁴ Furthermore, steatohepatitis caused by alcohol, drugs, or other forms of nonalcoholic steatohepatitis (NASH) share many histological features.⁵ The question is whether these disparate causes could lead to steatohepatitis and its potential sequelae of cirrhosis or liver failure by one or more common mechanisms. Any satisfactory unifying mechanism should ideally explain why, in some individuals, steatosis, whatever its etiology, never progresses to steatohepatitis,⁶ and also explain the variable incidence and severity of steatohepatitis and fibrosis in fatty liver of different etiologies. In this issue, Benson et al., working in Pessayre's laboratory, provide persuasive evidence that at least one such mechanism linking steatosis to necroinflammation and fibrosis is lipid peroxidation.⁷

A growing body of evidence supports a role for lipid peroxidation in the pathogenesis of alcohol-induced hepatitis and fibrosis.^{8,9} Ethanol metabolism results in the formation of reactive oxygen species (ROS) and carbon-centered free radicals capable of initiating peroxidation of the polyunsaturated fatty acid side chains of membrane phospholipids and lipoproteins. Potential sources of free radicals are the ethanol-inducible cytochrome P450 2E1 (CYP2E1), which generates superoxide, hydroxyl, and hydroxyethyl radicals, the mitochondrial respiratory chain (in response to the increased rate of reduced to oxidized nicotinamide adenine dinucleotide [NADH/NAD]), xanthine and aldehyde oxidases, and peroxisomal β -oxidation of free fatty acids, which generates hydrogen peroxide.¹⁰ CYP2E1-mediated generation of hydroxyethyl radicals in particular correlates closely with the degree of lipid peroxidation and liver damage in animal models of alcoholic liver disease (ALD).¹¹ A role for lipid peroxidation in NASH has been suggested by recent studies showing its presence in both animal models of nonalcoholic fatty liver and humans with steatosis of different etiologies.¹²⁻¹⁴

A major attraction of lipid peroxidation as an important common pathogenic mechanism of steatohepatitis associated with fatty liver of different etiologies is that it potentially explains most, if not all, of the diverse histological features observed in this condition. Peroxidation of membrane lipids may cause cell necrosis and megamitochondria. The aldehyde products of lipid peroxidation, 4-hydroxynonenal and malondialdehyde (MDA), are capable of activating hepatic stellate cells,^{15,16} the principal collagen-producing cells within the liver, cross-linking cytokeratins to form Mallory bodies,¹² and stimulating neutrophil chemotaxis.¹⁷ MDA may also contribute to inflammation by activating NF- κ B,¹⁸ a transcription factor regulating the expression of several proinflammatory cytokines and adhesion molecules including tumor necrosis factor α , interleukin 8, intercellular adhesion molecule 1, and E-selectin.¹⁹

Previous work from Pessayre's group has suggested that the mere presence of oxidizable fat within the liver is enough to trigger lipid peroxidation.¹² However, many patients with steatosis never progress to necroinflammation or fibrosis.⁶ This suggests that, in addition to steatosis (the first "hit"), the development of steatohepatitis requires the presence of some other factor(s) (second "hit"). This factor(s) might be expected to be particularly apparent in conditions where steatohepatitis is most commonly associated with steatosis, including alcohol-related liver disease and certain drugs. The results presented by Berson et al. show elegantly that one such second hit is a source of free radicals capable of inducing oxidative stress.

Many drugs are associated with the development of steatosis, including antiviral agents (interferon, zidovudine), aspirin, and other nonsteroidal anti-inflammatory drugs, the antiepileptic sodium valproate, and the tetracycline group of antibiotics. The fat is typically microvesicular in distribution and results predominantly from the inhibition of mitochondrial β -oxidation of fatty acids by a variety of different mechanisms.³ However, only one class of drugs is commonly associated with classical steatohepatitis, the cationic amphiphilic amines: amiodarone, perhexiline, and the coronary dilator 4,4'-diethylaminoethoxyhexestrol (DEAEH).^{3,20,21} Pessayre's group has shown previously that amiodarone^{22,23} and perhexiline²⁴ accumulate in the mitochondria and inhibit not only β -oxidation (causing steatosis, the first hit) but also the transfer of electrons along the respiratory chain. Theoretically, this latter effect would be expected to generate superoxide anions capable of initiating lipid peroxidation,²⁵ thus providing a potential mechanism for the second hit required for steatohepatitis to develop. In

this latest study they first show that, like amiodarone and perhexiline, DEAEH inhibits both β -oxidation and respiration, and then go on to show that all three drugs increase the production of ROS by isolated mitochondria and increase in vivo lipid peroxidation 5–10-fold. These data strongly suggest that it is because these drugs not only cause steatosis but also provide a mechanism for increased oxidative stress that they are capable of inducing steatohepatitis-like lesions.

This concept of steatohepatitis as a double-hit lesion potentially explains its relatively common occurrence in alcohol-related liver disease, where there exist several mechanisms for both steatosis¹ and oxidative stress. This theory may also explain why it has been difficult to develop animal models of alcoholic hepatitis and cirrhosis. In addition to simple ethanol administration, necroinflammation and fibrosis have only been produced by manipulations that have provided an alternative or extra source of oxidative stress. These have included increasing the dietary content of polyunsaturated fat (which induces the activity of CYP2E1²⁶) and iron²⁷ (which favors the production of hydroxyl radicals from hydrogen peroxide) and coadministering carbon tetrachloride vapor.²⁸ Because most heavy drinkers develop steatosis, the apparent individual susceptibility to advanced ALD seems likely to be explained by interindividual differences in the magnitude of the second hit, oxidative stress. This may be genetically determined, e.g., possession of the more transcriptionally active c2 allele of CYP2E1,²⁹ or influenced by environmental factors such as dietary intake of anti- or pro-oxidants.

The work of Pessayre's group might also be extrapolated to steatohepatitis of non-drug-related or ethanol-related etiologies. In particular, the data might explain why NASH occurs more commonly in association with some causes of steatosis than others and with greater severity. Some degree of lipid peroxidation can be shown in steatosis of most etiologies.¹² This presumably reflects the mild level of oxidative stress arising from normal physiological processes and is insufficient to cause significant liver injury. By analogy with drug-induced steatohepatitis, for NASH to occur, some additional source of oxidative stress (the second hit) is required that is capable of initiating enough lipid peroxidation to overcome the normal cellular defense mechanisms and produce necroinflammation. Recent studies have suggested several potential sources for this second hit. Increased expression of CYP2E1 has been shown in patients and animal models of NASH.^{30,31} In the absence of ethanol, CYP2E1 can generate free radicals from endogenously produced ketones and aldehydes and dietary *N*-nitrosamines. Possible

mediators of its induction in nonalcoholics include ketones and/or fatty acids,³² both of which may explain the induction of CYP2E1 by a high-fat diet.²⁶ Interestingly, among patients with obesity-related steatosis, the risk factors for steatohepatitis/fibrosis include rapid weight loss caused by dieting, debilitation or intestinal bypass surgery, surgical stress, alcohol intake, and diabetes, all of which are associated with an increase in the concentration of fatty acids and/or ketones within the liver.¹ In addition to CYP2E1 induction, an increase in the intrahepatic concentration of free fatty acids may provide a further source of oxidative stress via peroxisomal β -oxidation. This pathway becomes important in conditions of substrate overload or when mitochondrial β -oxidation is inhibited.³³ Unlike its mitochondrial counterpart, peroxisomal β -oxidation produces hydrogen peroxide that, in the presence of free iron, is converted to the highly reactive hydroxyl radical. The importance of liver iron in disease pathogenesis has been illustrated by a recent study showing that patients with NASH have an increased frequency of the C282Y mutation in the recently cloned hemochromatosis gene, HFE.³⁴ Clearly, as for ALD, other genetic determinants of oxidative stress could also play a role in susceptibility to NASH.

This study by Berson et al. of a small group of drugs has provided a basis for understanding at least one of the important mechanisms of steatohepatitis regardless of etiology. Its development requires a double hit, the first producing steatosis, the second a source of oxidative stress capable of initiating significant lipid peroxidation. This concept provides a rationale for both the treatment and prevention of disease progression in steatosis of alcoholic and nonalcoholic causes.

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